

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 10/591,726
Source: IFWP
Date Processed by STIC: 9/13/06

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 09/13/2006

PATENT APPLICATION: US/10/591,726

TIME: 11:07:38

Input Set : A:\21085064U1.TXT

Output Set: N:\CRF4\09132006\J591726.raw

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4 <110> APPLICANT: UAB Research Foundation
6 <120> TITLE OF INVENTION: BRHF1 AS A CANCER DIAGNOSTIC MARKER
9 <130> FILE REFERENCE: 21085.0064P1
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/591,726
C--> 11 <141> CURRENT FILING DATE: 2006-09-05
11 <150> PRIOR APPLICATION NUMBER: 60/550,224
12 <151> PRIOR FILING DATE: 2004-03-04
14 <160> NUMBER OF SEQ ID NOS: 21
16 <170> SOFTWARE: FastSEQ for Windows Version 4.0
18 <210> SEQ ID NO: 1
19 <211> LENGTH: 105
20 <212> TYPE: PRT
21 <213> ORGANISM: Artificial Sequence
23 <220> FEATURE:
24 <223> OTHER INFORMATION: Description of Artificial Sequence:/note =
25     Synthetic Construct
27 <400> SEQUENCE: 1
28 Met Lys Gly Leu Ser Pro Ile Ala Lys Gly Arg Lys Thr Ser Val Ser
29 1           5           10           15
30 Ala Ala Val Leu Val Ser Thr Thr Ile Pro Ile Ser Ser Val Trp Gly
31           20           25           30
32 Pro Leu Gln Ile Leu Gly Gln Lys Arg Gly Gln Lys Met Glu Gln Ala
33           35           40           45
34 Asn His Pro Val Gly Leu Asp Ile Ser Val Val Tyr Lys Asp Thr Leu
35           50           55           60
36 Lys Lys Ile Val Gln Gln Glu Thr Ser Cys Pro Phe Thr His Val His
37 65           70           75           80
38 Tyr Ala Glu Gly Ile Thr Gly Arg His Thr Ala Pro Glu Asp Glu Gly
39           85           90           95
40 Ser Leu Ala Gln Lys Pro Pro Ile Arg
41           100          105
44 <210> SEQ ID NO: 2
45 <211> LENGTH: 270
46 <212> TYPE: PRT
47 <213> ORGANISM: Artificial Sequence
49 <220> FEATURE:
50 <223> OTHER INFORMATION: Description of Artificial Sequence:/note =
51     Synthetic Construct
53 <400> SEQUENCE: 2
54 Met Asn Ile Asp Ala Lys Ile Leu Asn Lys Ile Leu Ala Asn Gln Ile
55 1           5           10           15
56 Gln Gln His Ile Lys Lys Leu Ile His His Asp Gln Val Gly Phe Ile
57           20           25           30

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58 Pro Gly Met Gln Gly Trp Phe Asn Ile His Lys Ser Ile Asn Val Ile
59          35          40          45
60 Gln His Ile Asn Arg Thr Lys Asp Lys Asn His Met Ile Ile Ser Val
61          50          55          60
62 Asp Ala Glu Lys Ala Phe Asp Lys Val Gln Gln His Phe Met Leu Lys
63 65          70          75          80
64 Thr Leu Asn Lys Leu Gly Ile Asp Gly Thr Tyr Leu Lys Ile Ile Arg
65          85          90          95
66 Ala Ile Tyr Asp Lys Pro Thr Ala Asn Ile Ile Leu Asn Gly Leu Lys
67          100          105          110
68 Leu Glu Ala Phe Pro Leu Lys Thr Gly Thr Arg Gln Gly Cys Pro Leu
69          115          120          125
70 Ser Leu Leu Leu Phe Asn Ile Val Leu Glu Val Leu Ala Arg Ala Ile
71          130          135          140
72 Arg Gln Glu Lys Glu Ile Asn Cys Ile Gln Leu Gly Lys Glu Glu Val
73 145          150          155          160
74 Lys Leu Pro Leu Phe Ala Asp Asp Met Ile Val Tyr Leu Glu Asn Pro
75          165          170          175
76 Val Val Ser Ala Pro Asn Leu Leu Lys Leu Ile Ser Asn Phe Ser Lys
77          180          185          190
78 Val Ser Gly Tyr Lys Ile Asn Val Gln Lys Ser Gln Ala Phe Leu Tyr
79          195          200          205
80 Thr Asn Asn Arg Gln Thr Glu Ser Gln Ile Met Ser Glu Leu Pro Phe
81          210          215          220
82 Thr Ile Ala Ser Lys Arg Ile Lys Tyr Leu Gly Ile Gln Leu Thr Arg
83 225          230          235          240
84 Asp Val Lys Asp Leu Phe Lys Glu Asn Tyr Lys Pro Leu Leu Asn Glu
85          245          250          255
86 Ile Lys Glu Asp Thr Asn Lys Cys Lys Asn Ile Pro Cys Ser
87          260          265          270
90 <210> SEQ ID NO: 3
91 <211> LENGTH: 315
92 <212> TYPE: DNA
93 <213> ORGANISM: Artificial Sequence
95 <220> FEATURE:
96 <223> OTHER INFORMATION: Description of Artificial Sequence:/note =
97     Synthetic Construct
99 <400> SEQUENCE: 3
100 atgaagggat tatcgctat cgccaagggg aggaaaacta gtgtttctgc tgctgtgttg 60
101 gtgagcacia ctattccgat cagcagtgtc tggggaccat tgcagattct tgggcaaaag 120
102 agaggacaga aaatggagca ggccaatcac ccagtggggc ttgatatcag tgtgggttac 180
103 aaggacacct taaaaaagat tgtccaacaa gaaacaagct gccccttcac ccatgtccac 240
104 tatgctgagg gaatcactgg aaggcacact gcccagagg atgaaggttc tctggcccag 300
105 aagcccccaa tcaga                                     315
107 <210> SEQ ID NO: 4
108 <211> LENGTH: 810
109 <212> TYPE: DNA
110 <213> ORGANISM: Artificial Sequence
112 <220> FEATURE:

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113 <223> OTHER INFORMATION: Description of Artificial Sequence:/note =

114 Synthetic Construct

116 <400> SEQUENCE: 4

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117 atgaacatcg atgcaaaaat cctcaataaa atactggcaa accaaatcca gcagcacatc 60
118 aaaaagctta tccaccatga tcaagtgggc ttcacccctg ggatgcaagg ctggttcaac 120
119 atacacaaat caataaatgt aatccagcat ataaacagaa ccaaagacaa aaaccacatg 180
120 attatctcag tagatgcaga aaaggccttt gacaaagttc aacaacactt catgctaaaa 240
121 actctcaata aattaggtat tgatgggacg tatctcaaaa taataagagc tatctatgac 300
122 aaaccacacag ccaatatcat actgaatggg ctaaaactgg aagcattccc ttgaaaact 360
123 ggcacaagac agggatgccc tctctcactt ctcctattca acatagtgtt ggaagttctg 420
124 gccagggcaa tcaggcagga gaaggaaata aattgtattc aattaggaaa agaggaagtt 480
125 aaattgcccc tgtttgacga tgacatgatt gtatatctgg aaaaccccg tctctcagcc 540
126 ccaaattctcc ttaagctgat aagcaacttc agcaaagtct caggatacaa aatcaacgtg 600
127 caaaaatcac aagcattctt atacaccaat aacagacaaa cagagagcca aatcatgagt 660
128 gaactcccat tcacaattgc ttcaaagaga ataaaatacc taggaatcca acttacaagg 720
129 gatgtgaagg acctcttcaa ggagaactac aaaccactgc tcaacgaaat aaaagaggat 780
130 acaaacaaat gcaagaacat tccatgctca 810

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132 <210> SEQ ID NO: 5

133 <211> LENGTH: 1263

134 <212> TYPE: DNA

135 <213> ORGANISM: Artificial Sequence

137 <220> FEATURE:

138 <223> OTHER INFORMATION: Description of Artificial Sequence:/note =

139 Synthetic Construct

141 <400> SEQUENCE: 5

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142 catctacaga actctccacc ccaaataaac agaataataca tttttttcag caccacacca 60
143 cacctattcc aaaattgacc acatagtgtg aagtaaagct ctcctcagca aatgtaaaag 120
144 aacagaaatt ataacaaact atctctcaga ccacagtgca atcaaactag aactcaggat 180
145 taagaatctc actcaaagcc gctcaactac atggaaactg aacaacctgc tcctgaatga 240
146 ctactgggta cataacgaaa tgaaggcaga aataaagatg ttctttgaaa ccaacgagaa 300
147 caaagacacc acatacaga atctctggga cgcattcaaa gcagtgtgta gagggaaatt 360
148 tatagacta aatgcctacc agagaaagca ggaagatcc aaaattgaca ccctaacatc 420
149 acaattaaaa gaactagaaa agcaagagca aacacattca aaagctagca gaaggcaaga 480
150 aataactaaa atcagagcag aactgaagga aatagagaca caaaaaacc ttcaaaaaat 540
151 caatgaatcc aggagctggt tttttgaaag gatcaacaaa attgatagac cgctagcaag 600
152 actaataaag aaaaaaagag agaagaatca aatagacaca ataaaaaatg ataaagggga 660
153 tatcaccacc gatccacag aaatacaaac taccatcaga gaatactaca aacacctcta 720
154 cgcaataaaa ctagaaaatc tggaagaaat ggatacattc ctcgacacat acactctccc 780
155 aagactaaac caggaagaag ttgaatctct gaatcgacca ataacaggct ctgaaattgt 840
156 ggcaataatc aatagtttac caacaaaaa gagtccagga ccagatggat tcacagccga 900
157 attctaccag aggtacaagg aggaactggt accattcctt ctgaaactat tccaatcaat 960
158 agaaaaagag ggaatcctcc ctaactcatt ttatgagacc agcatcattc tgataccaaa 1020
159 gccgggcaga gacacaacca aaaaagagaa ttttagacca atatccttga tgaacattga 1080
160 tgcaaaaatc ctcaataaaa tactggcaaa ccgaatccag cagcacatca aaaagcttat 1140
161 ccaccatgat caagtgggct tcatccctgg gatgcaaggc tggttcaata tacgcaaatc 1200
162 aataaatgta atccagcata taaacagagc caaagacaaa aaccacatga ttatctcaat 1260
163 aga 1263

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165 <210> SEQ ID NO: 6

166 <211> LENGTH: 10

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167 <212> TYPE: DNA
168 <213> ORGANISM: Artificial Sequence
170 <220> FEATURE:
171 <223> OTHER INFORMATION: Description of Artificial Sequence:/note =
172     Synthetic Construct
174 <400> SEQUENCE: 6
175 cagagcctgt                                     10
177 <210> SEQ ID NO: 7
178 <211> LENGTH: 10
179 <212> TYPE: DNA
180 <213> ORGANISM: Artificial Sequence
182 <220> FEATURE:
183 <223> OTHER INFORMATION: Description of Artificial Sequence:/note =
184     Synthetic Construct
186 <400> SEQUENCE: 7
187 ctctgggaca                                     10
189 <210> SEQ ID NO: 8
190 <211> LENGTH: 375
191 <212> TYPE: PRT
192 <213> ORGANISM: Artificial Sequence
194 <220> FEATURE:
195 <223> OTHER INFORMATION: Description of Artificial Sequence:/note =
196     Synthetic Construct
198 <400> SEQUENCE: 8
199 Met Lys Gly Leu Ser Pro Ile Ala Lys Gly Arg Lys Thr Ser Val Ser
200 1          5          10          15
201 Ala Ala Val Leu Val Ser Thr Thr Ile Pro Ile Ser Ser Val Trp Gly
202          20          25          30
203 Pro Leu Gln Ile Leu Gly Gln Lys Arg Gly Gln Lys Met Glu Gln Ala
204          35          40          45
205 Asn His Pro Val Gly Leu Asp Ile Ser Val Val Tyr Lys Asp Thr Leu
206          50          55          60
207 Lys Lys Ile Val Gln Gln Glu Thr Ser Cys Pro Phe Thr His Val His
208 65          70          75          80
209 Tyr Ala Glu Gly Ile Thr Gly Arg His Thr Ala Pro Glu Asp Glu Gly
210          85          90          95
211 Ser Leu Ala Gln Lys Pro Pro Ile Arg Met Asn Ile Asp Ala Lys Ile
212          100         105         110
213 Leu Asn Lys Ile Leu Ala Asn Gln Ile Gln Gln His Ile Lys Lys Leu
214          115         120         125
215 Ile His His Asp Gln Val Gly Phe Ile Pro Gly Met Gln Gly Trp Phe
216          130         135         140
217 Asn Ile His Lys Ser Ile Asn Val Ile Gln His Ile Asn Arg Thr Lys
218 145         150         155         160
219 Asp Lys Asn His Met Ile Ile Ser Val Asp Ala Glu Lys Ala Phe Asp
220          165         170         175
221 Lys Val Gln Gln His Phe Met Leu Lys Thr Leu Asn Lys Leu Gly Ile
222          180         185         190
223 Asp Gly Thr Tyr Leu Lys Ile Ile Arg Ala Ile Tyr Asp Lys Pro Thr

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224      195      200      205
225 Ala Asn Ile Ile Leu Asn Gly Leu Lys Leu Glu Ala Phe Pro Leu Lys
226      210      215      220
227 Thr Gly Thr Arg Gln Gly Cys Pro Leu Ser Leu Leu Leu Phe Asn Ile
228 225      230      235      240
229 Val Leu Glu Val Leu Ala Arg Ala Ile Arg Gln Glu Lys Glu Ile Asn
230      245      250      255
231 Cys Ile Gln Leu Gly Lys Glu Glu Val Lys Leu Pro Leu Phe Ala Asp
232      260      265      270
233 Asp Met Ile Val Tyr Leu Glu Asn Pro Val Val Ser Ala Pro Asn Leu
234      275      280      285
235 Leu Lys Leu Ile Ser Asn Phe Ser Lys Val Ser Gly Tyr Lys Ile Asn
236      290      295      300
237 Val Gln Lys Ser Gln Ala Phe Leu Tyr Thr Asn Asn Arg Gln Thr Glu
238 305      310      315      320
239 Ser Gln Ile Met Ser Glu Leu Pro Phe Thr Ile Ala Ser Lys Arg Ile
240      325      330      335
241 Lys Tyr Leu Gly Ile Gln Leu Thr Arg Asp Val Lys Asp Leu Phe Lys
242      340      345      350
243 Glu Asn Tyr Lys Pro Leu Leu Asn Glu Ile Lys Glu Asp Thr Asn Lys
244      355      360      365
245 Cys Lys Asn Ile Pro Cys Ser
246      370      375
249 <210> SEQ ID NO: 9
250 <211> LENGTH: 1125
251 <212> TYPE: DNA
252 <213> ORGANISM: Artificial Sequence
254 <220> FEATURE:
255 <223> OTHER INFORMATION: Description of Artificial Sequence:/note =
256     Synthetic Construct
258 <400> SEQUENCE: 9
259 atgaagggat tatcgcttat cgccaagggg agggaaaacta gtgtttctgc tgctgtgttg      60
260 gtgagcacia ctattccgat cagcagtgtc tggggaccat tgcagattct tgggcaaaag      120
261 agaggacaga aaatggagca ggccaatcac ccagtggggc ttgatatcag tgtgggtttac      180
262 aaggacacct taaaaaagat tgtccaacaa gaaacaagct gccccttcac ccatgtccac      240
263 tatgtctgagg gaatcactgg aaggcacact gcccagagg atgaagggtt tctggccag      300
264 aagcccccaa tcagaatgaa catcgatgca aaaatcctca ataaaatact ggcaaaccac      360
265 atccagcagc acatcaaaaa gcttatccac catgatcaag tgggcttcat ccctgggatg      420
266 caaggctggt tcaacataca caaatcaata aatgtaatcc agcatataaa cagaacccaa      480
267 gacaaaaacc acatgattat ctcatgatg gcagaaaagg cctttgacaa agttcaacaa      540
268 cacttcatgc taaaaactct caataaatta ggtattgatg ggacgtatct caaaataata      600
269 agagctatct atgacaaacc cacagccaat atcatactga atgggctaaa actggaagca      660
270 ttccctttga aaactggcac aagacaggga tgccctctct cacttctcct attcaacata      720
271 gtgttggaag ttctggccag ggcaatcagg caggagaagg aaataaattg tattcaatta      780
272 ggaaaagagg aagttaaatt gcccctgttt gcagatgaca tgattgtata tctggaaaac      840
273 cccgtcgtct cagccccaaa tctccttaag ctgataagca acttcagcaa agtctcagga      900
274 tacaaaatca acgtgcaaaa atcacaagca ttcttatata ccaataacag acaaacagag      960
275 agccaaatca tgagtgaact ccatttcaca attgcttcaa agagaataaa atacctagga      1020
276 atccaactta caagggatgt gaaggacctc ttcaaggaga actacaaacc actgctcaac      1080

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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/591,726

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Input Set : A:\21085064U1.TXT

Output Set: N:\CRF4\09132006\J591726.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date